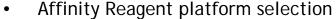
Preliminary Conclusions

- Target Selection
 - Literature based?
 - Expression based?
 - Mass Spectrometry based?
 - Proposal based?



- Are antibodies really the best?
- Monoclonal vs. polyclonal?
- What about other scaffolds/platforms?
- What should we use as the determinants?
 - Experience
 - Available secondary reagents
 - Intellectual property consideration
- Should we pick a platform or just let everyone play?

Antigens

- Should these be produced by each affinity reagent producer?
- Should these be made as a centralized resource?

Target Selection

- Depends largely on the project's overall goals
- User requested:
 - Focused collections of proteins, networks, pathways of interest to motivated individuals in the community:
 - Many of these pathways have proven themselves to be critical to disease and possibly biomarkers
 - Guarantees that there will be users who will employ antibodies

But...

- There are already mechanisms in place to obtain antibodies for these proteins
- Project Design:
 - Centralized process (could involve as many sources as needed) to select antigens for a specific purpose
 - Acknowledges the goal to make affinity reagents for <u>infrequently</u> studied proteins
 - Could be based on:
 - Literature mining several sources
 - Specific data that indicate likely candidates for specified purpose
 - Abundance data e.g., look at proteins that are rare in serum
 - Predicted properties of proteins potential solubility, extra-cellular domains, likely to be secreted
 - Avoid duplication with existing antibodies!

Preliminary Conclusions

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Reagent Platform

- Major bottlenecks are reagent platform independent:
 - Making antigen
 - Validating affinity reagent
- Antibodies are still the most mature technology – tens of thousands already available
- Alternative platforms show promise but still need further development
- Monoclonals cost more because individual clones must be screened, but have the advantage that they are a renewable resource
 - This is important because once the reagent is validated it becomes much more valuable
- Polyclonals cost less and could be very powerful for screening candidates
- Working with a consistent platform would simplify high throughput (proteome scale) applications
- Current producers can make between 500 - 1500 monoclonals per year

Preliminary Conclusions

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Antigens

- Making antigens is a key bottleneck
- They are useful in both production and validation phases
- Affinity reagent producers would like a central source for antigen
- Success rates for affinity reagents depend heavily on the ability to make good quality antigen
- Not clear how to organize and manage a centralized source for antigen

Preliminary Conclusions II

- Mechanisms for contracting the production of the reagents
 - Should we worry about avoiding duplicating effort?
 - OR Duplication can be good and done at the risk of the producer
 - Should we pay only for finished antibodies?
 - Must meet established criteria
 - Validated for pre-selected applications



- Once acquired, the target gets crossed off the list
- Should we contract each producer in advance to make a target list of antibodies?
 - Milestones must be met to complete payment
- Should this effort focus primarily on a centralized mechanism for validating affinity reagents and storing this information?
 - Would a "seal of approval" and the opportunity to list the reagent in this "central database" be an incentive to develop these reagents?

Production Process

- Validation is the most important part of this process.
- Creating a database that tracks various reagents, lists their qualities, and "certifies" them somehow is highly desirable (more details below)
- Commercial mechanisms are in place to produce antibodies to <u>popular</u> proteins - companies are happy to take these suggestions
 - Some will partner with academics to make these antibodies at low or no cost
- Incentives are needed to get producers to create antibodies to <u>infrequently</u> studied proteins.
 - Paying for production would ensure that NCI owned the antibodies
 - Paying for validation i.e., providing it via central source - might induce production of some antibodies
 - "Suggesting proteins" might stimulate some production (but if the proteins are not frequently studied - this may not get traction)

Preliminary Conclusions III

Distribution

- Option 1: Centralized repository/distribution center
- Option 2: Centralized database/producers handle their own distribution through standard commercial mechanisms



- Both? Other?
- IP considerations for distribution - limits?
- If affinity reagents are contracted and paid for by NIH, who owns the IP rights to the reagents?

Distribution

- Centralized Distribution has the advantages that:
 - Antibody production and distribution can be handled with consistent standards regarding QC/QA, antibody concentration, etc.
 - Having all the affinity reagents in one location will simplify the development of high throughput applications
 - Simplify the MTA and IP morass
- Centralized distribution has the disadvantages that:
 - Requires infrastructure
 - Duplicates an existing network
 - Realistically could only apply to reagents produced and paid for by a centralized effort.

Preliminary Conclusions IV

- Validation
 - Which assays should be validated for?
 - Denatured protein
 - Western blot
 - ELISA
 - Immunohistochemistry
 - Native protein
 - Immunoprecipitation
 - Antibody arrays
 - Other...



- How many affinity reagents needed per target?
- Which criteria must be met to accept an antibody?
 - E.g., if an antibody is very good at only one of the above, is this ok?
- Should validation be done at centralized facilities?
 - Or is it ok for various centers to validate using the same criteria?

Validation

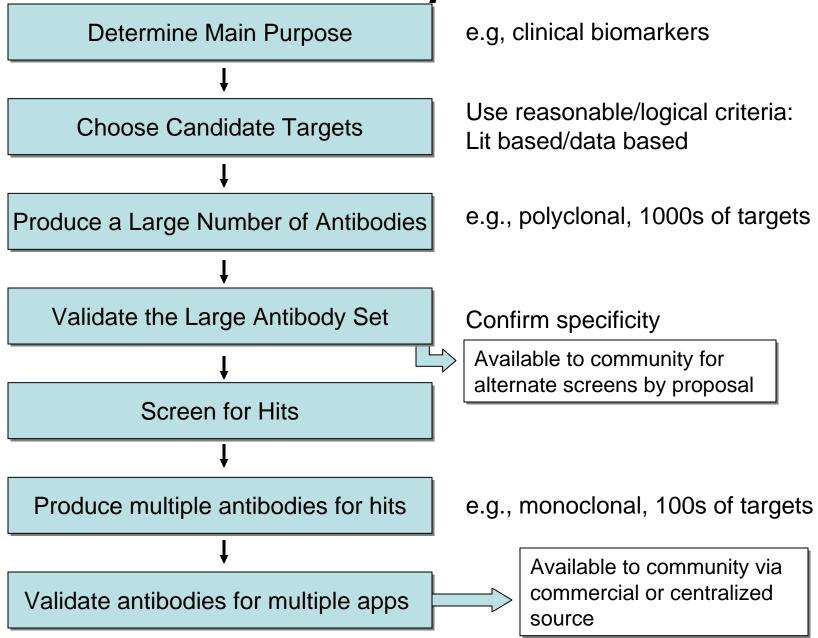
- Strong support for a centralized database that lists the characteristics and qualities for all antibodies, including existing ones
 - Data from existing antibodies could be provided voluntarily by labs using them
 - Specific criteria/format vs. at the labs' own discretion
- A centralized validation process is needed for reagents generated under a planned project.
 - Consistent standards and SOPs are applied here
 - Data is centralized and qualifying reagents could be "certified"
 - Would need to periodically revisit all reagents to monitor QA/QC
- Applications to validate include:
 - Standard applications like westerns, ELISA, etc. - usually done as part of antibody development
 - Applications that foster high throughput uses

Straw II

Proposal

- Select a single key motivation for developing the reagents
 - i.e., clinical biomarkers
- Select targets based on reasonable criteria
 - Literature/biology based
 - Existing relevant data
- Produce a large number of candidate antibodies to the targets, e.g., polyclonal
 - Available to community on a proposal basis
- Centralized core validation confirms specificity
- Stage 1: Screen all the antibodies for the key application
- Stage 2: Take all the "hits" and produce multiple antibodies to each, e.g., monoclonal
- Both centralized and distributed validation of monoclonals
 - Available to community via central or commercial network

Straw Proposal II



Straw II

Assumptions

- It's expensive to make and validate multiple antibodies to each protein - this limits the number of antigens that can be targeted
- Target selection depends on the proposed use for the targets
- A main thrust of this project is to develop reagents to proteins not frequently studied already
 - There are already strong commercial motivations to make antibodies to <u>frequently studied proteins</u>
- At least one main thrust of this project is to find new clinical biomarkers for detecting/categorizing disease from serum

Straw II

Advantages

- Focuses on proteins not currently studied
- Many more first stage antibodies can be made
- Opportunity to screen for many "hits"
- Monoclonals only invoked on "interesting proteins"
- Reduces money spent on less interesting proteins

Disadvantages

- Focuses on only one main application
- Polyclonal antibodies not renewable
- Proteins that are not hits in this assay might be important in another